

FIG. - 1

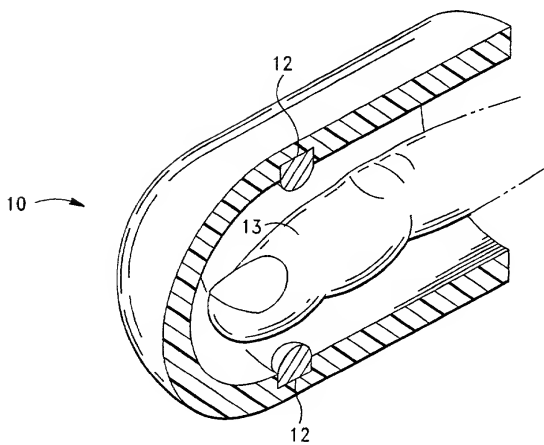
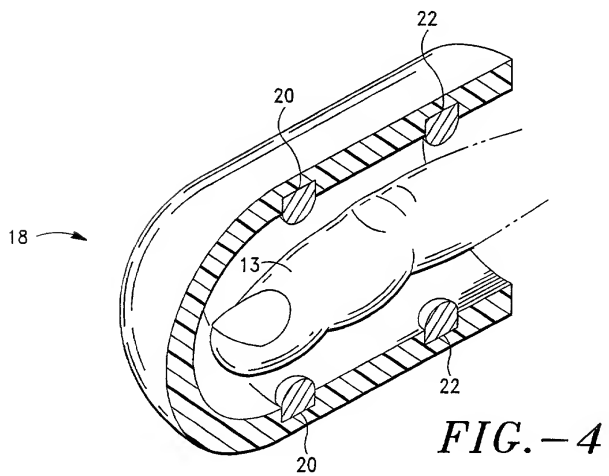
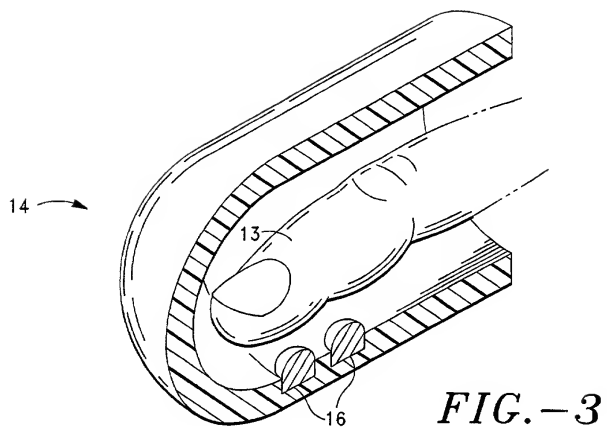


FIG. - 2



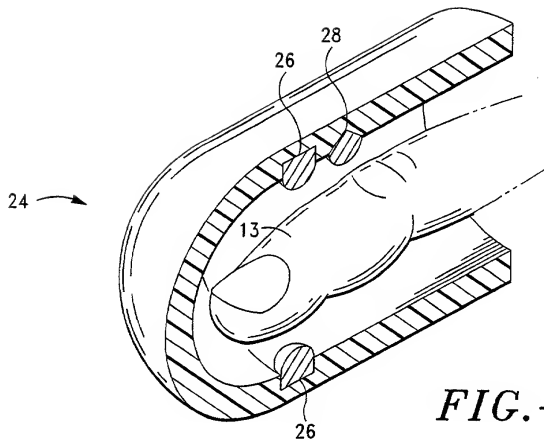


FIG. -5

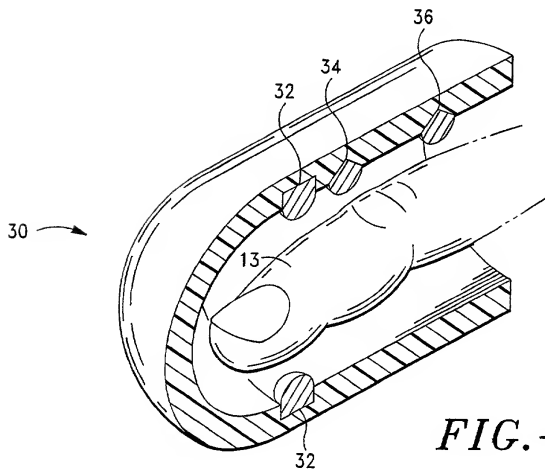


FIG. -6

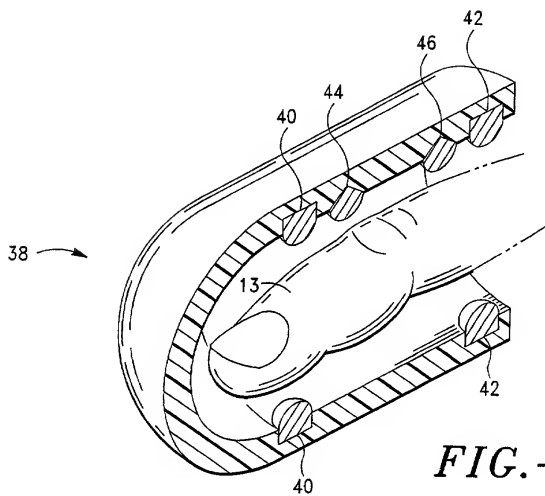


FIG. - 7

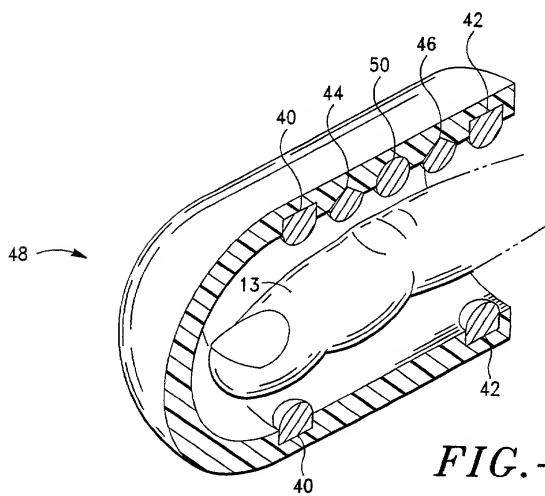


FIG. - 8

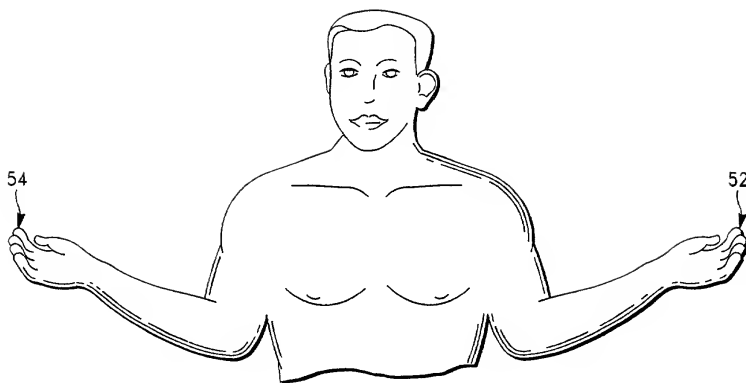


FIG. - 9

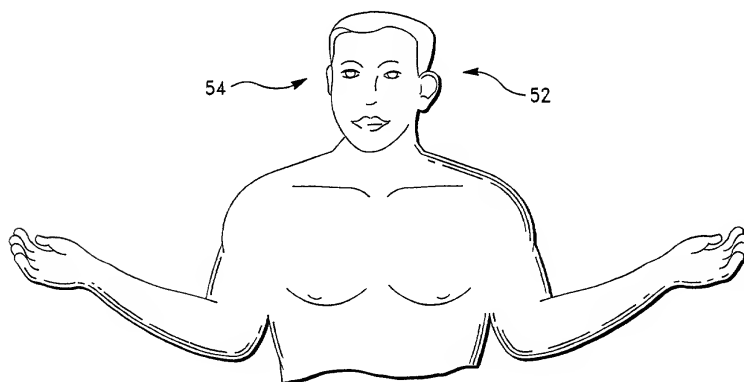


FIG. - 10

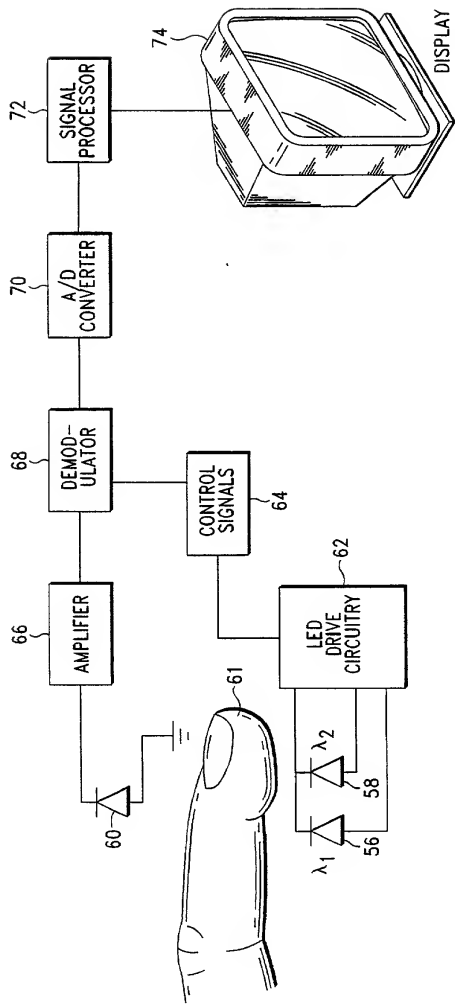


FIG.-11

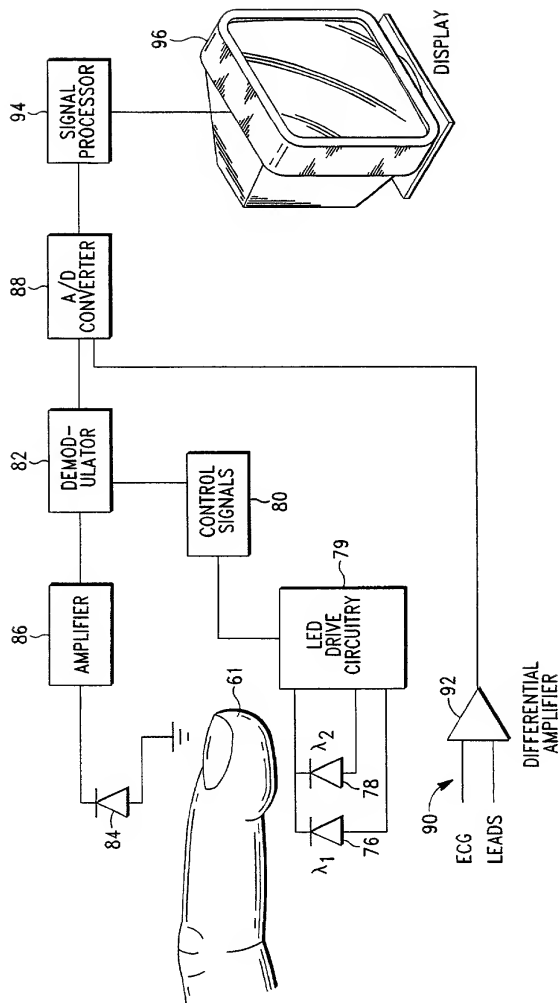
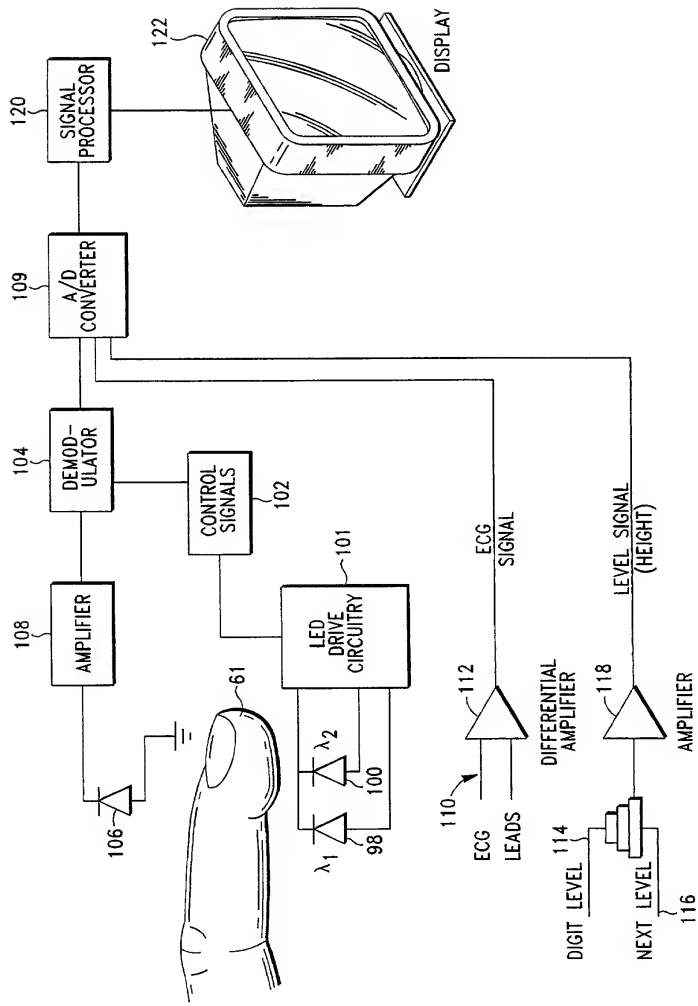


FIG.-12



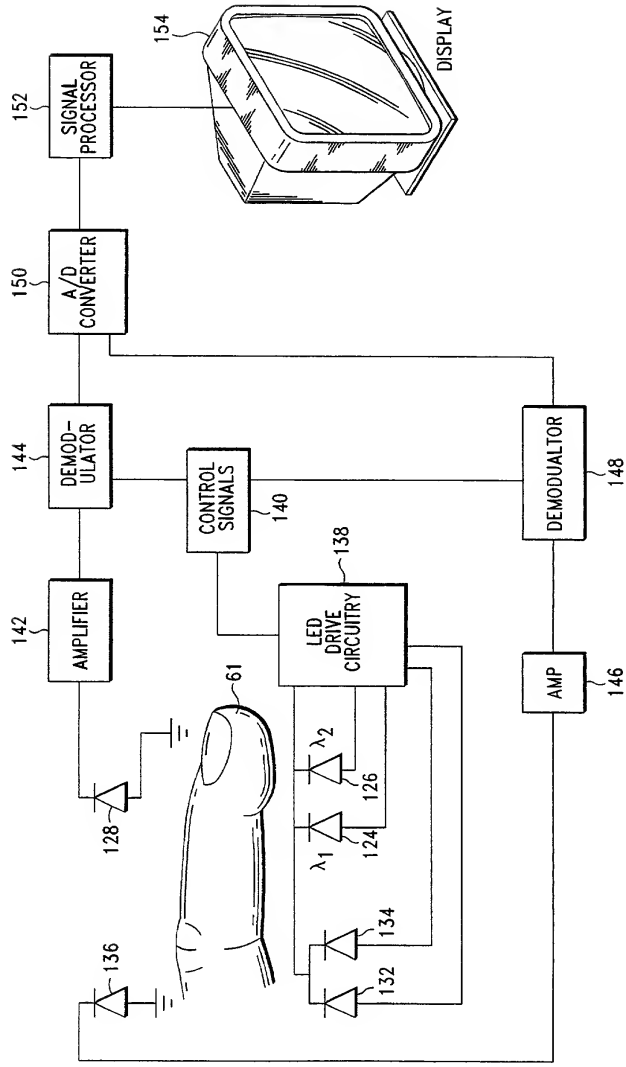


FIG.-14

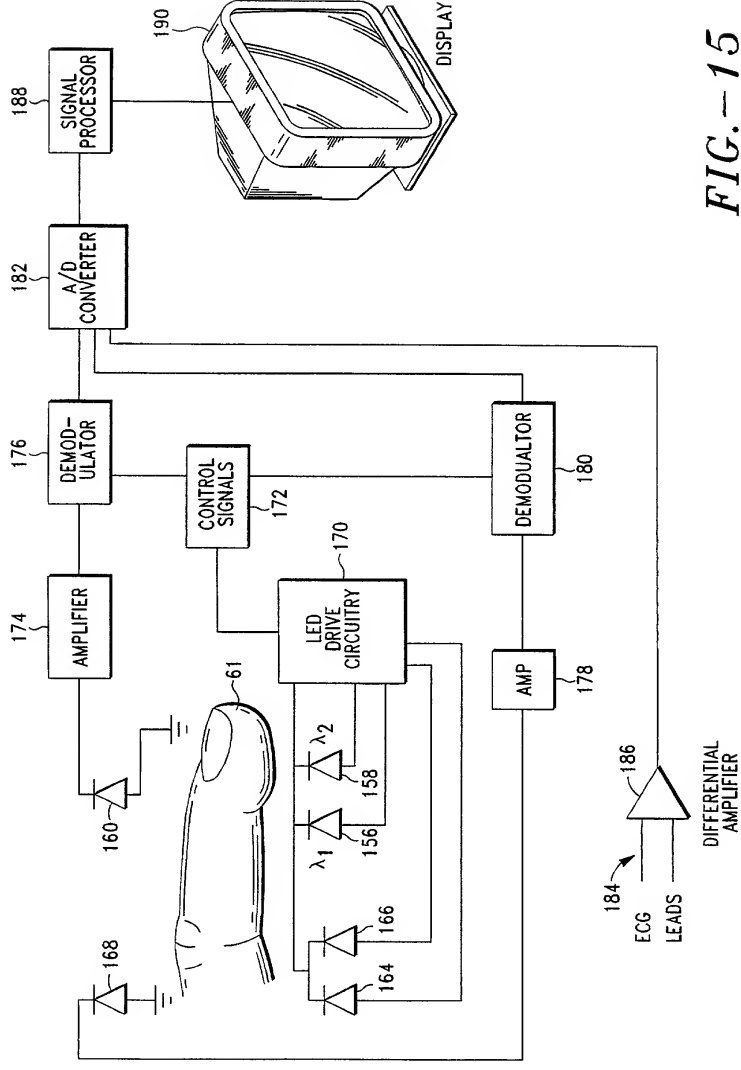


FIG.-15

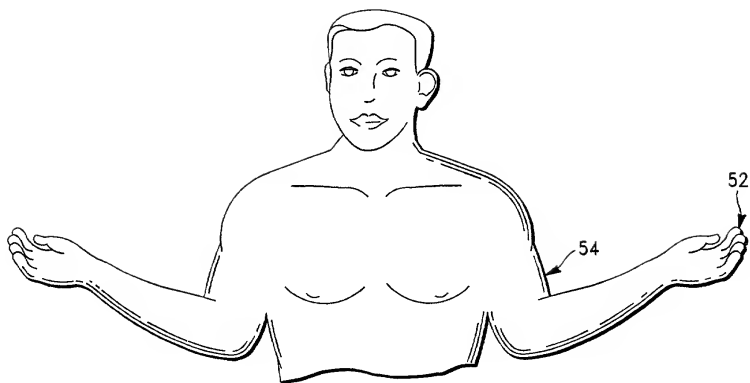


FIG. - 16

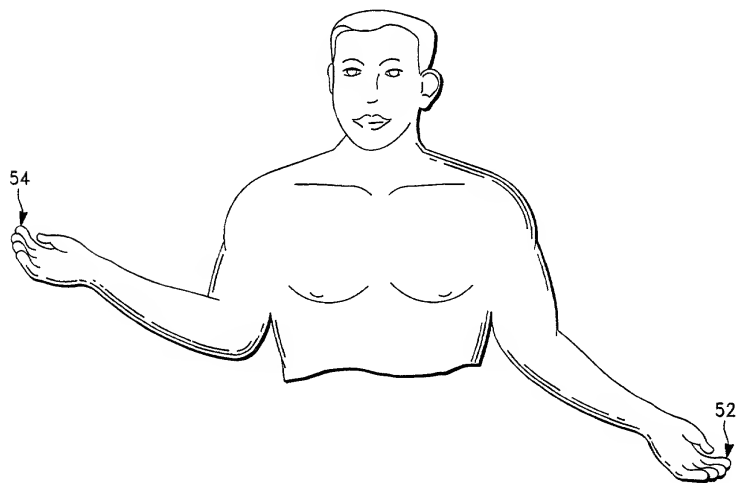


FIG. - 19

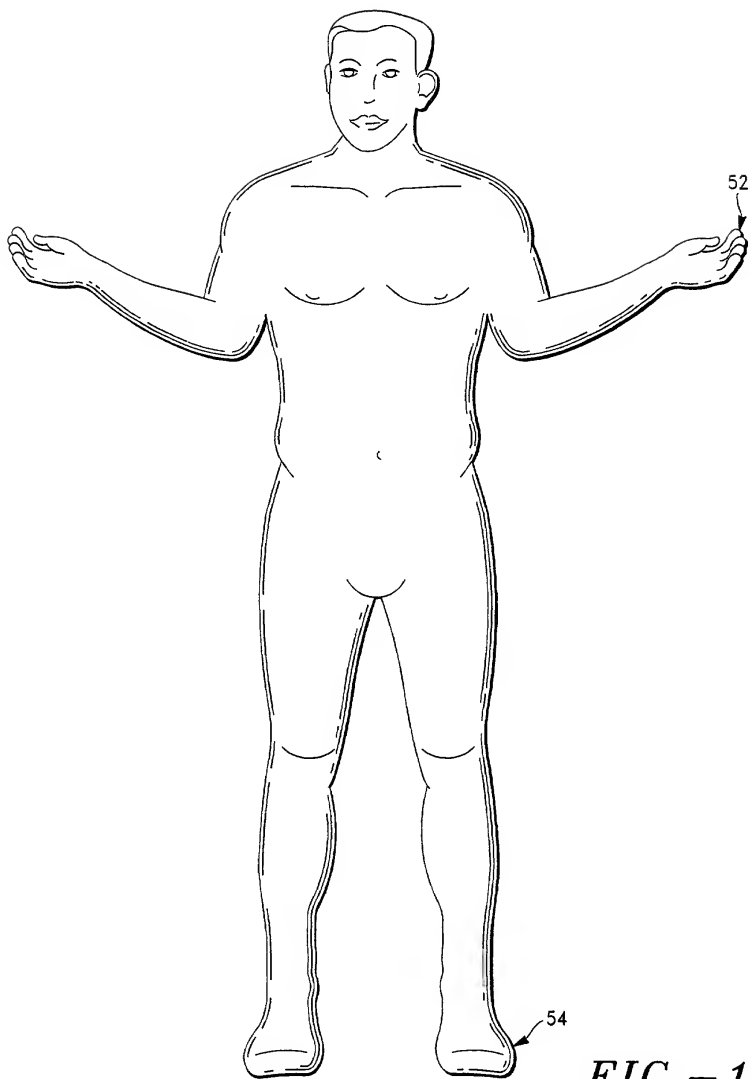


FIG.-17

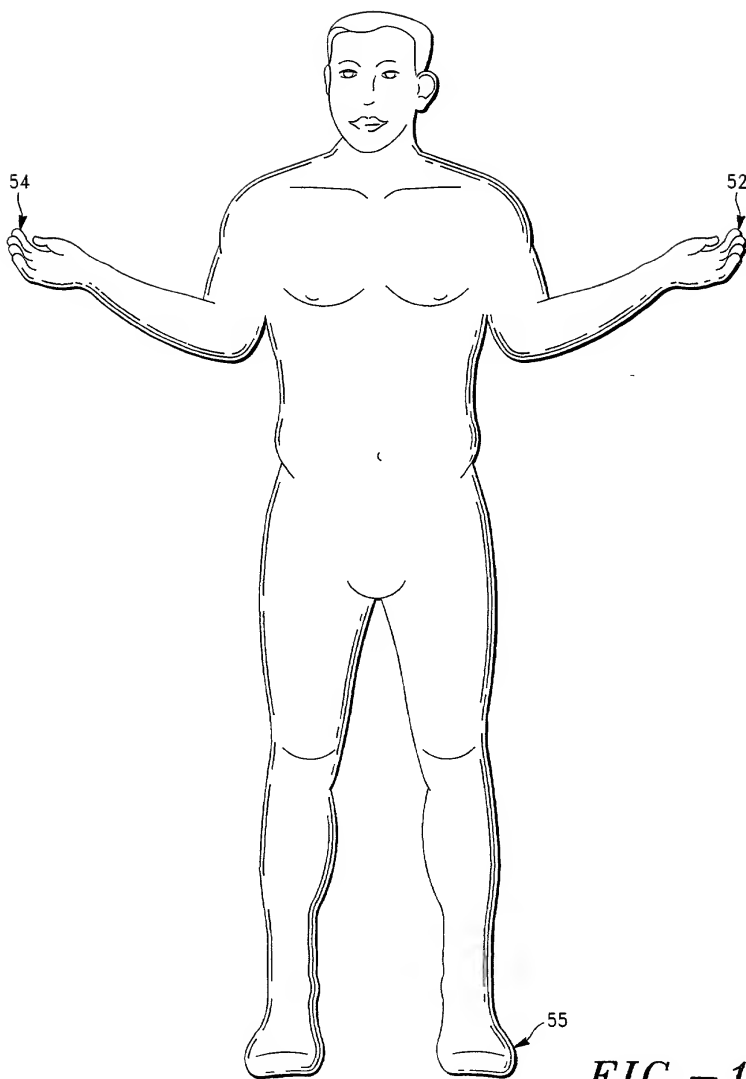


FIG.-18

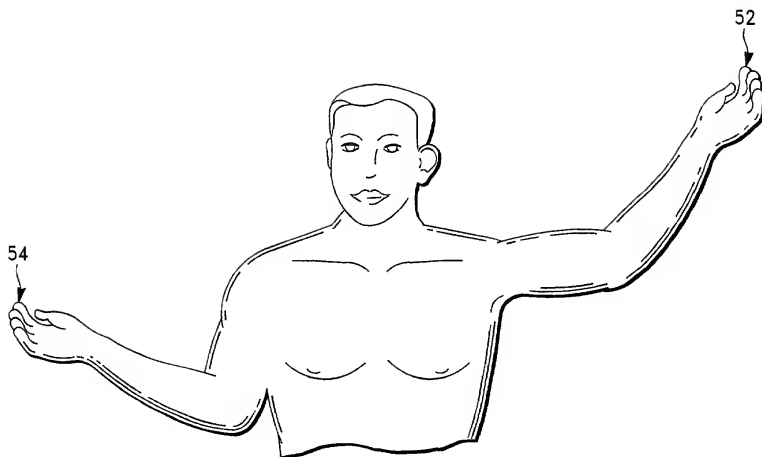


FIG.—20

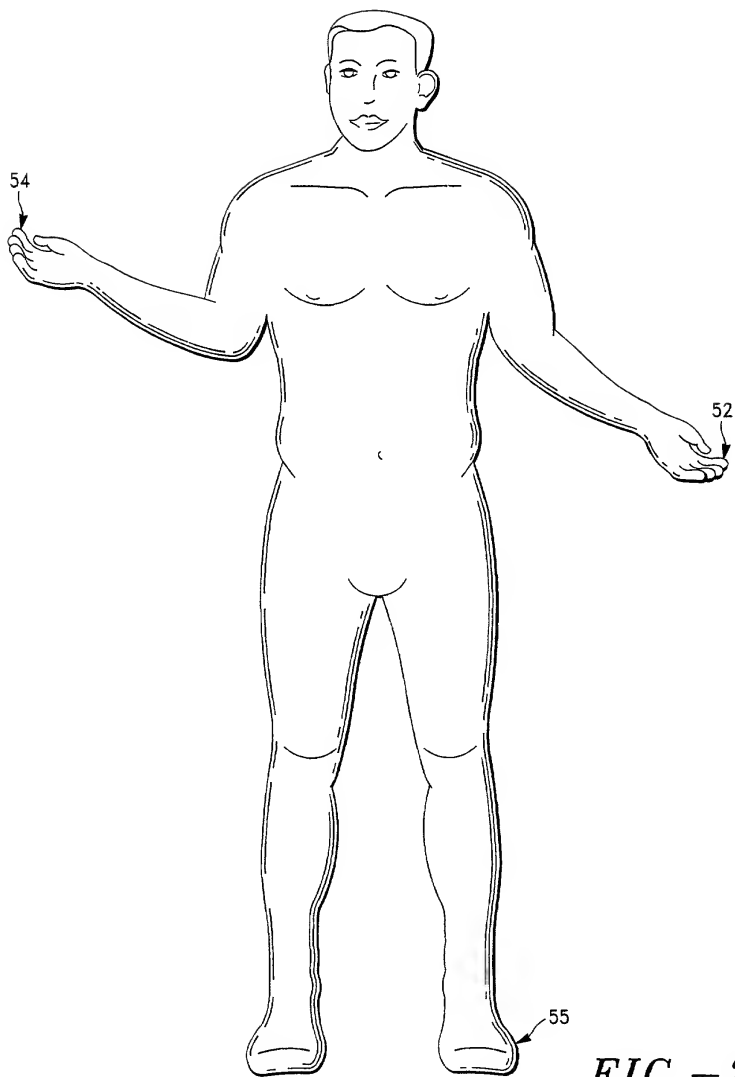


FIG.-21

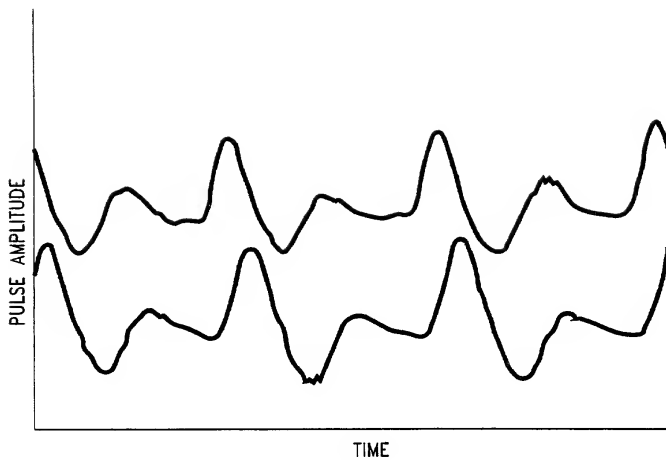


FIG.-22

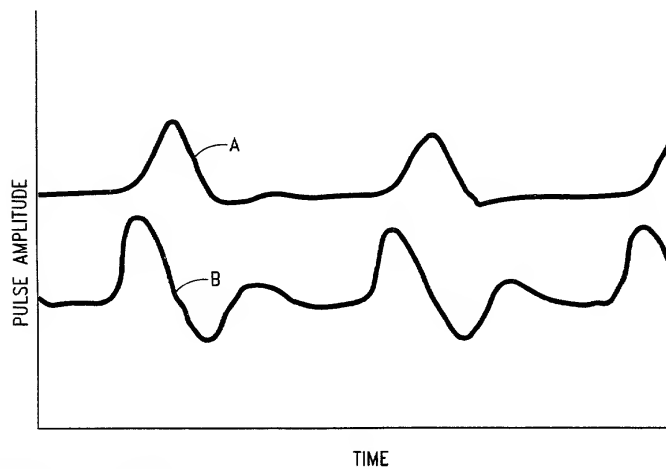


FIG.-23

The graph plots Pulse Amplitude on the vertical axis against Time on the horizontal axis. Two waveforms are shown: A (upper) and B (lower). Both waveforms exhibit a periodic, irregular pattern. Signal A has a higher amplitude and more pronounced peaks and troughs compared to signal B, which has a lower amplitude and a flatter profile. The two signals are in phase, meaning their peaks and troughs occur at approximately the same time intervals.

The graph displays two pulse waveforms, A and B, plotted against Time. The vertical axis is labeled 'PULSE AMPLITUDE' and the horizontal axis is labeled 'TIME'. Waveform A is a square wave with sharp transitions, while Waveform B is a smooth, periodic wave.

FIG.-25

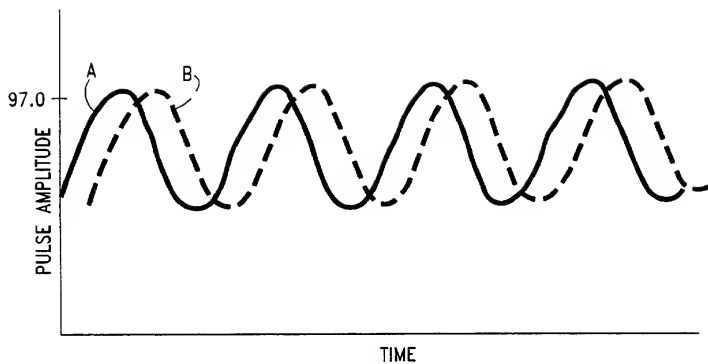


FIG. - 26

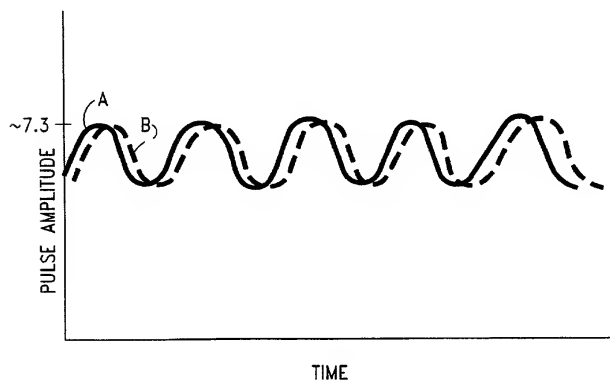


FIG. - 27

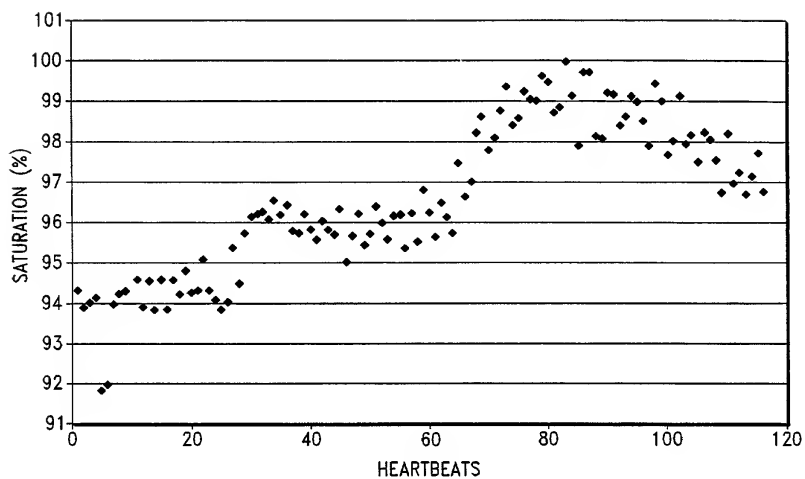


FIG.-28

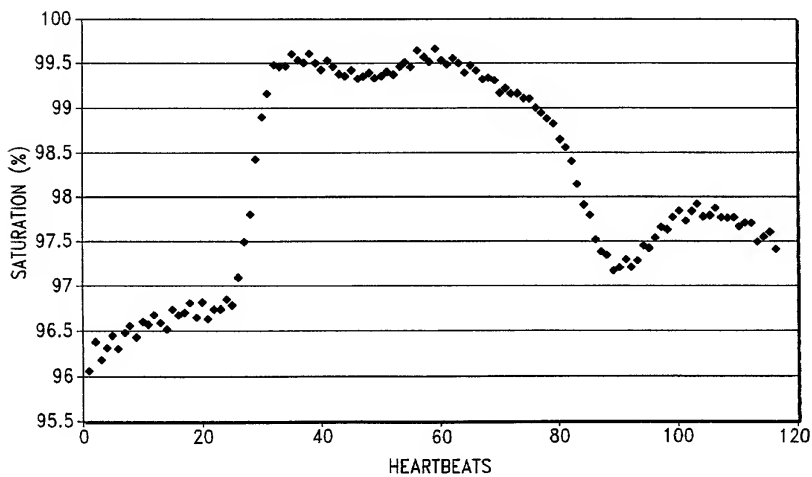


FIG.-29

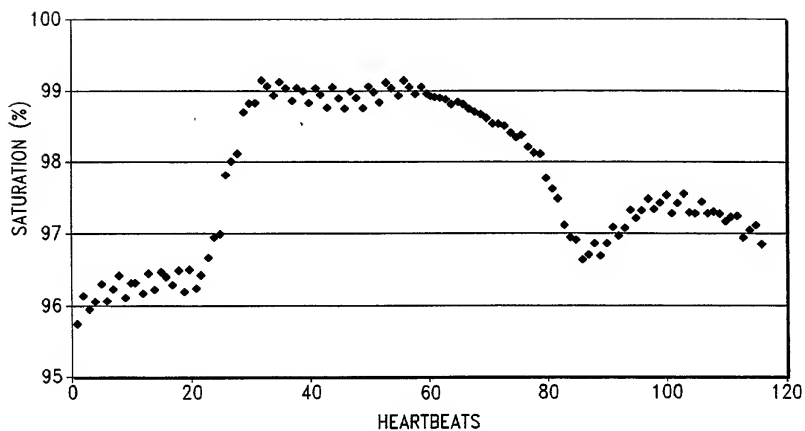


FIG.-30

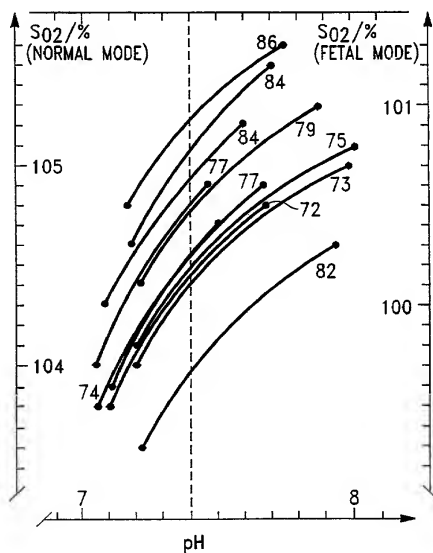


FIG.-31

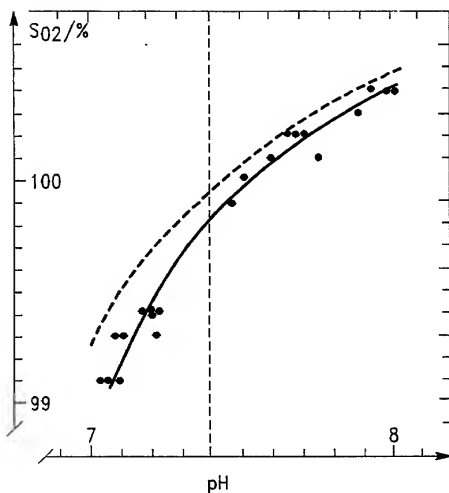


FIG.-32

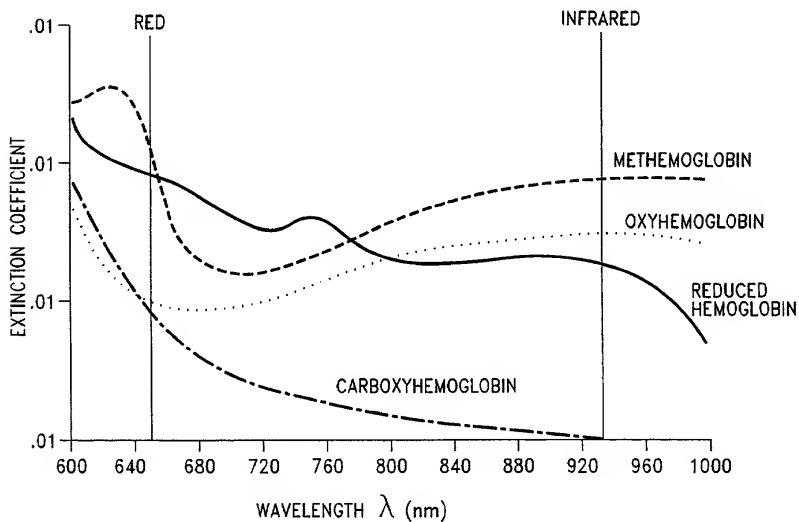


FIG.-34

Computational Algorithm for Determination of Hemoglobin Concentration

```

C      A is the measured absorbance
C      A1 is the absorbance after dividing out extinction coefficients
C      and correcting for saturation
C      A2,A3, ... will be the absorbances at different path lengths,
C      created by multiplying by constants
C      A1,A3, ... and L2,L3, etc.
C      constant M2=0.9
C      constant M3=0.8
C      constant M4=0.7
C      constant M5=0.6
C      constant M6=0.5
C      constant M7=0.4
C      constant M8=0.3

C
C      read in the value for hemoglobin absorbance and a value k
C      representing the extinction coefficient for the wavelength and
C      the oxygen saturation
Begin
Read, A
Read, k
A1:=A/k
A2:=A1*M2
A3:=A1*M3
A4:=A1*M4
A5:=A1*M5
A6:=A1*M6
A7:=A1*M7
A8:=A1*M8

C
C      k1234 = log(A1) * log(A2) - log(A3) * log(A4)
C      k5678 + log(A5) * log(A6) - log(A7) * log(A8)
C      kd:=[ log(A1*A2) - log(A3*A4) ] / [ log(A5*A6) - log(A7*A8) ]

C
C      combine all the A terms that occur as coefficients.
C      kAc := log(A2/A1) - log(A3/A1) - log(A4/A1) - [(kd * log(A5/A1)) -
C      - [kd * log(A6/A1) + [kd * log(A7/A1)] + [kd * log(A8/A1)]

C
C      combine all the A terms that occur alone
C      kAa :=- [log(A3/A1) * log(A4/A1)] ) -
C      - kd * [log(A5/A1) * log(A6/A1)] +
C      + kd* [(log(A7/A1) * log(A8/A1)]

C
C      k1234 - ( kd * k5678) = kig(L) * kAc + kAa
C      log(L) = [k1234 - (kd * k5678) - kAa] / kAc
C      L = antilog{[k1234 - (kd * k5678) - kAa] / kAc}
C      use EXP or antilog function
C      L = EXP([k1234 - (kd * k5678) - kAa] / kAc)
C      L is the path length
C      C is the concentration of hemoglobin
C      C = A1 / L

C
END

```

FIG.—33

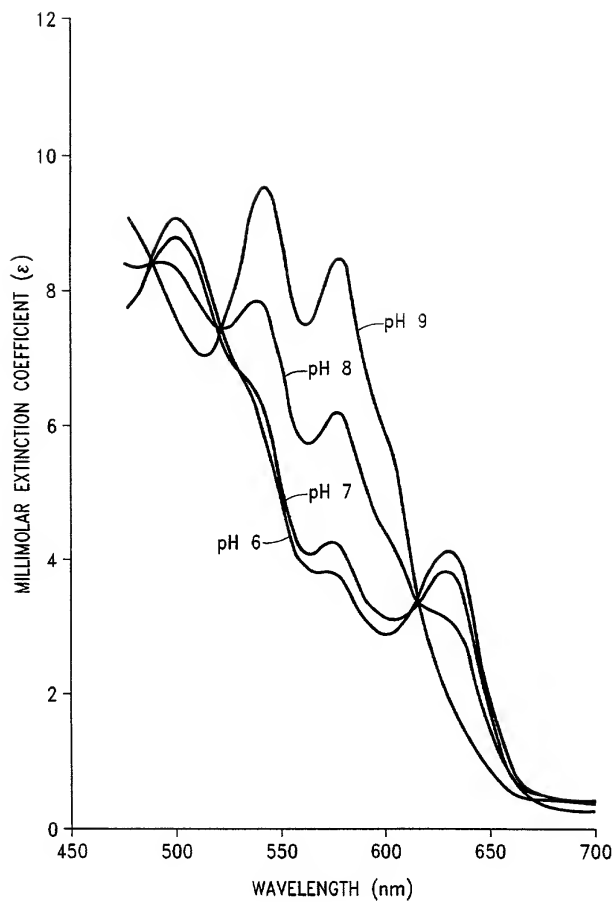


FIG. - 35

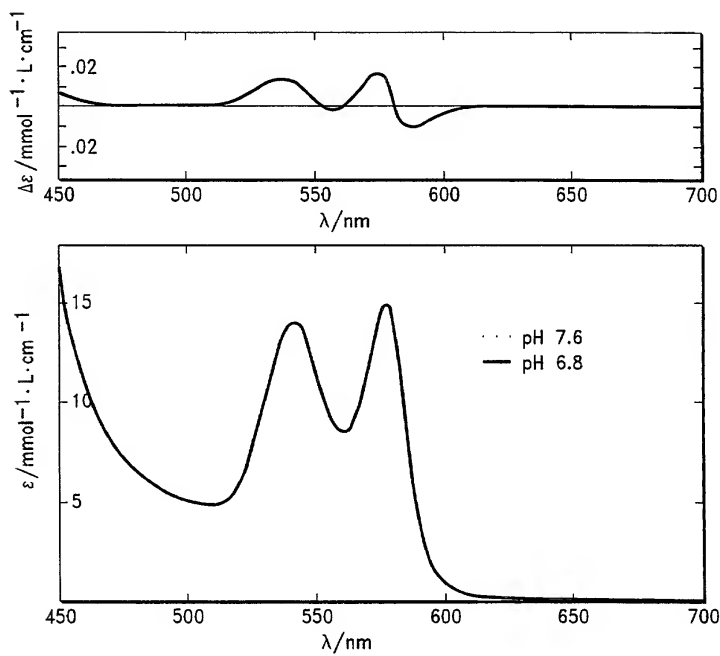


FIG. - 36

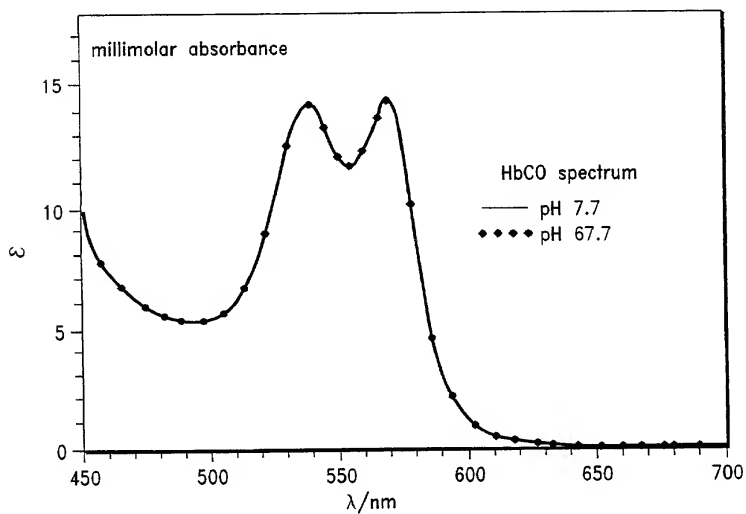
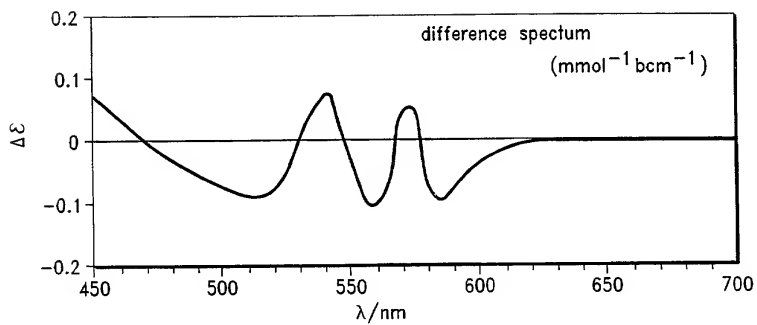


FIG. - 37

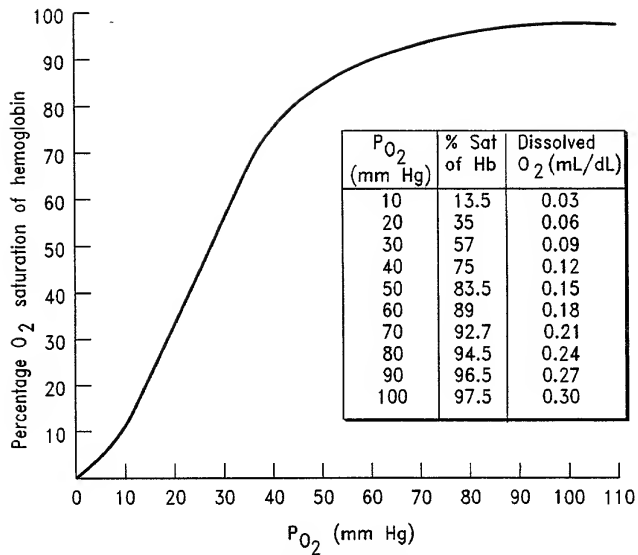


FIG.-38

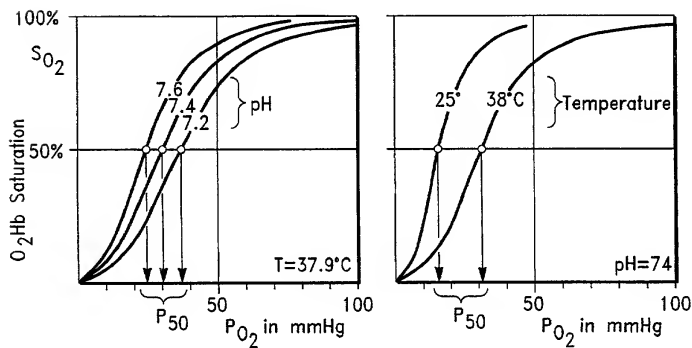


FIG.-39

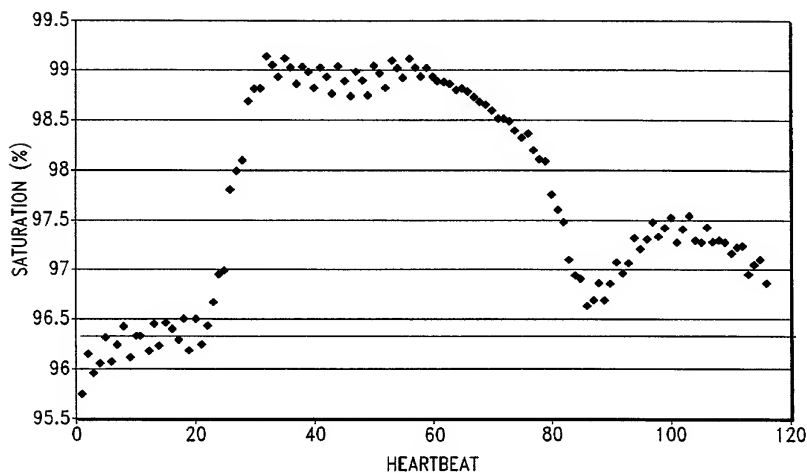


FIG.-40

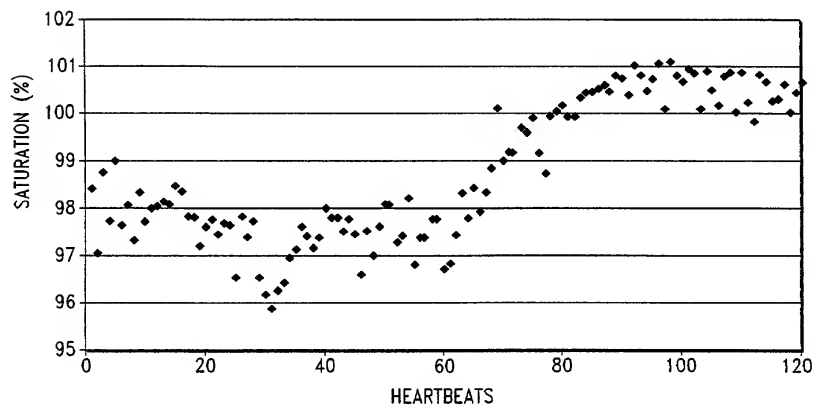


FIG.-41